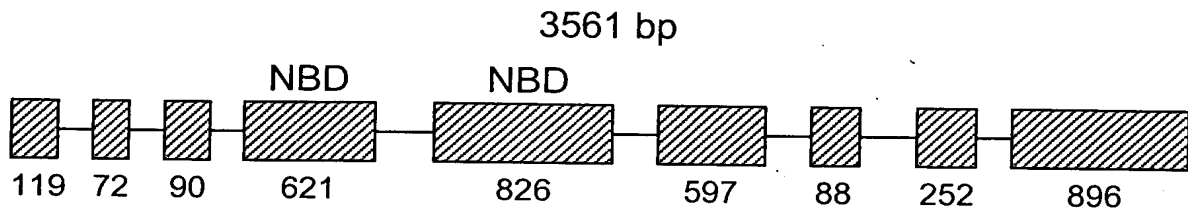
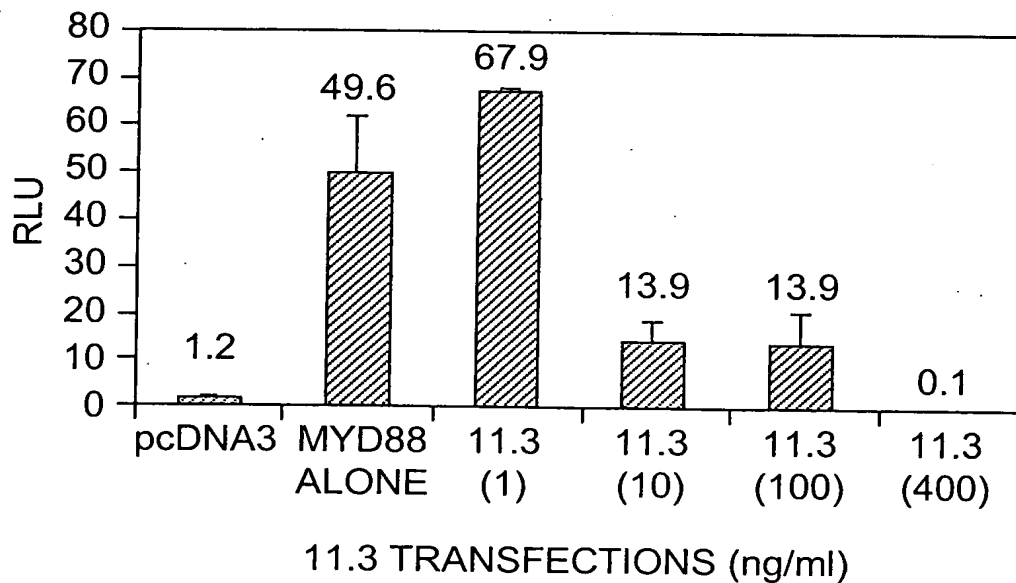


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101 GPTFALDTVHVDPIRESTPDELLRPPAELALEHQPPQAGLPPLALSOLF
151 NPDACGRRVQTVVLYGTGTGKSTLVRKMVLDCYGRLPAPFELLIPFSCE
201 DLSSLGPAPASLCQLVAQRYTPLKEVLPLMAAAGSHLLFVLHGLEHLNLD
251 FRLAGTGLCSDPEEPQEPAAIIVNLLRKYMLPQASILVTTRPSAIGRIPS
301 KYVGRYGEICGFSDTNLQKLYFQLRLNQPYCGYAVGGSGVSATPAQRDHL
351 VQMLSRNLEGGHQAIAACFLPSYCWLVCATLHFLHAPTAPAGQTLTSIYTS
401 FLRLNFSGETLDSTDPNLSLMAYAARTMGKLAYEGVSSRKTYFSEEDVC
451 GCLEAGIRTEEEFQLLHIFRRDALRFFLAPCVEPGRAGTFVFTVPAMQEQY
501 LAALYIVLGLRKTTLQKVGKEVAELVGRVGEDVSLVLGIMAKLLPLRALP
551 LLFNLIKVVPRVFGRMVGKSREAVTQAMVLEMFREEDYNDVLDQMGAS
601 ILGVEGPRRHPDEPPEDEVFELFPMFMGGLLSAHNRAVLAQLGCPIKNLD
651 ALENAQAIKKKLGLGRQVLPPSELDDHLFFHYEFQNRFSAEVLSSLRQ
701 LNLAGVRMTPVKCTVVAAVLGSGRHALDEVNLAQCQLDPAGLRTLLPVFL
751 RARKLGLQLNSLGPACDLDRLDLLHDQCQITTLRLSNPLTEAGVAVLM
801 EGLAGNTSVTHLSLLHTGLGDEGLELLAAQLDRNRQLQELNVAYNGAGDT
851 AALALARAAREHPSLELLQGVAIQMCWKLPLLPYAHLWTPRMPSHWCFL
901 ILMPLPQWYDGLVAPRGRCT (SEQ ID NO:20)

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FIG. 25G**FIG. 26****FIG. 27**
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